

# **INFORMATION DISCLOSURE STATEMENT BY APPLICANT** ( Not for submission under 37 CFR 1.99)

Application Number	10617979
Filing Date	2003-07-11
First Named Inventor	Henkin et al.
Art Unit	1637
Examiner Name	Samuel C. Woolwine
Attorney Docket Number	22727/04130

## **U.S. PATENTS**

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/S.W./	1	2004/007677	WO		2004-01-22	The Ohio State University		<input type="checkbox"/>

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/S.W./	1	Artsimovitch, I., et al., "RNA polymerases from <i>Bacillus subtilis</i> and <i>Escherichia coli</i> differ in recognition of regulatory signals in vitro", (2000) <i>J. Bacteriol.</i> 182, 6027–6035.	<input type="checkbox"/>
	2	Grandoni, J. A., et al., "Regions of the <i>Bacillus subtilis</i> <i>ilv-leu</i> Operon involved in regulation by Leucine" (1993) <i>J. Bacteriol.</i> 175, 7581–7593.	<input type="checkbox"/>
	3	Grundy, F. J., et al., "Interaction between the acceptor end of tRNA and the T box stimulates antitermination in the <i>Bacillus subtilis</i> <i>tyrS</i> gene: a new role for the discriminator base" (1994) <i>J. Bacteriol.</i> 176, 4518–4526.	<input type="checkbox"/>
	4	Grundy, F. J., et al., "tRNA determinants for transcription antitermination of the <i>Bacillus subtilis</i> <i>tyrS</i> gene". (2000) <i>RNA</i> 6, 1131–1141.	<input type="checkbox"/>
	5	Grundy et al., "Monitoring uncharged tRNA during transcription of the <i>Bacillus subtilis</i> <i>glyQS</i> Gene", (2005) <i>J. Mol. Biol.</i> 348, 73–81.	<input type="checkbox"/>
	6	Hager, D. A., et al., "Use of mono Q high-resolution ion-exchange chromatography to obtain highly pure and active <i>Escherichia coli</i> RNA polymerase", (1990) <i>Biochemistry</i> 29, 7890–7894.	<input type="checkbox"/>
	7	Hurwitz et al., "The intracellular concentration of bound and unbound magnesium ions in <i>Escherichia coli</i> ", (1967) <i>J. of Biol. Chemistry</i> , 242, 3719–3722.	<input type="checkbox"/>
	8	Landick, R., Turnbough, C. L., Jr., and Yanofsky, C. (1996) in <i>Escherichia coli</i> and <i>Salmonella</i> : Cellular and Molecular Biology, eds. Neidhardt, F. C., Curtis, R., III, Ingraham, J. L., Lin, E. C. C., Low, K. B., Magasanik, B., Reznikoff, W. S., Riley, M., Schaechter, A. and Umberger, H. E. (Am. Soc. Microbiol., Washington, DC), 1263–1286.	<input type="checkbox"/>
	9	Luo, D., et al., "In vitro and in vivo secondary structure probing of the <i>thrS</i> leader in <i>Bacillus subtilis</i> ", (1998) <i>Nucleic Acids Res.</i> 26, 5379–5387.	<input type="checkbox"/>
	10	Nelson et al., "tRNA regulation of gene expression: Interactions of an mRNA 5'-UTR with a regulatory tRNA", (2006) <i>RNA</i> , 12, 1–8.	<input type="checkbox"/>
↓	11	Qi, Y. & Hulett, F. M. "PhoP~P and RNA polymerase $\sigma^A$ holoenzyme are sufficient for transcription of <i>Pho</i> regulon promoters in <i>Bacillus subtilis</i> : PhoP~P activator sites within the coding region stimulate transcription in vitro", (1998) <i>Mol. Microbiol.</i> 28, 1187–1197.	<input type="checkbox"/>

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/S.W./	12	Rollins, S. M., et al., "Analysis of cis-acting sequence and structural elements required for antitermination of the bacillus subtilis tyrS gene", (1997) Mol. Microbiol. 25, 411-421.	<input type="checkbox"/>
	13	Winkler, W. C., et al., "The GA motif: an RNA element common to bacterial antitermination systems, rRNA, and eukaryotic RNAs", (2001) RNA 7, 1165-1172.	<input type="checkbox"/>
	14	Yousef et al., "Structural transitions induced by the interaction between tRNAGLY and the bacillus subtilis glyQS T box leader RNA", (2005) J Mol Biol, 349, 273-287.	<input type="checkbox"/>
	15	Henkin et al., "Sensing Metabolic Signals with nascent RNA transcripts: the T-box and S-box riboswitches as paradigms", (2007) Cold Spring Harbor Symposia on Quantitative Biology, vol. LXXI, 1-7.	<input type="checkbox"/>
	16	Grundy, F. J. & Henkin, T. M. "tRNA as a positive regulator of transcription antitermination in B. subtilis", (1993) Cell 74, 475-482.	<input type="checkbox"/>
	17	Anagnostopoulos, C. & Spizizen, J. "Requirements for Transformation in Bacillus Subtilis", (1961) J. Bacteriol. 81, 741-746.	<input type="checkbox"/>
	18	Ban et al., "The Complete Atomic Structure of the Large Ribosomal Subunit at 2.4 Å Resolution", (2000) Science 289, 905-920.	<input type="checkbox"/>
	19	Friedman, D. I. & Court, D. L. "Bacteriophage lambda: alive and well and still doing its thing", (2001) Curr. Opin. Microbiol. 4, 201-207.	<input type="checkbox"/>
	20	Giege et al., "Universal rules and idiosyncratic features in tRNA identity", (1996) Nucleic Acids Res. 26, 5017-5035.	<input type="checkbox"/>
	21	Grundy et al., "Regulation of the Bacillus subtilis Acetate Kinase Gene by CcpA", (1993) J. Bacteriol. 175, 7348-7355.	<input type="checkbox"/>
↓	22	Ogle et al., "Recognition of Cognate Transfer RNA by the 30S Ribosomal Subunit", (2001) Science 292, 897-902.	<input type="checkbox"/>

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/S.W./	23	Qiu et al., "The tRNA-binding moiety in GCN2 contains a dimerization domain that interacts with the kinase domain and is required for tRNA binding and kinase activation", (2001) EMBO J. 20, 1425-1438.	<input type="checkbox"/>
	24	Rhodes, G. & Chamberlin, M. J. "Ribonucleic Acid Cain Elongation by Escherichia coli Ribonucleic Acid Polymerase", (1974) J. Biol. Chem. 249, 6675-6683.	<input type="checkbox"/>
	25	Sankaranarayanan et al., "The Structure of Threonyl-tRNA Synthetase-tRNA Complex Enlightens Its Repressor Activity and Reveals an Essential Zinc Ion in the Active Site", (1999) Cell 97, 371-381.	<input type="checkbox"/>
	26	Treilber, D. K. & Williamson, J. R. "Beyond kinetic traps in RNA folding", 82, 221-230.(2001) Curr. Opin. Struct. Biol. 11, 309-314.	<input type="checkbox"/>
	27	Weeks, K. M. & Cech, T. R. "Protein Facilitation of Group I Intron Splicing by Assembly of the Catalytic Core and the 5' Splice Site Domain", (1995) Cell 82, 221-230.	<input type="checkbox"/>
↓	28	Guerrier-Takada et al., "The RNA Moiety of Ribonuclease P is the Catalytic Subunit of the Enzyme", (1983) Cell 35, 849-857.	<input type="checkbox"/>

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Examiner Signature	/Samuel Woolwine/	Date Considered	05/05/2008
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